

Appl. No. 08/935,377; Filed: September 22, 1997
 Dkt. No. 1821.0010000/EKS/HCC; Group Art Unit: 1644
 Inventors: Maurice ZAUDERER; Tel: 202/371-2600
 Title: Methods for Selecting Polynucleotides Encoding T Cell
 Epitopes (as amended)



p7.5tk

7.5 k promoter

Not I Apa I

GGCCAATTGAAACTAGCTTATTGCAC**GGGGGGCGATG** GGC CTC G GGC GGC AAC GGC GGA
 Met Gly Pro Ala Ala Asn Gly Gly

tk coding sequence

pE/L tk

synthetic E/L promoter

Not I Apa I

GGCCAATTGAAATTATTTGGATATAAG**GGGGCGCTATG** GGC CTC G GGC AAC GGC GGA
 Met Gly Pro Ala Ala Asn Gly Gly

tk coding sequence

FIG. 1

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1. p7.5tk

7.5K PROMOTER

NOTI

APAI

5'- GGCCAAAAATTGAAAAACTAGATCTATTATTGCACGGGCCGCATGGGGCCGCC-3'

2. p7.5/ATG0/tk

7.5K PROMOTER

NOTI

BAMHI SMAI PSTI

5'- GGCCAAAAATTGAAAAACTAGATCTATTATTGCACGGGCCGCATGGGGCCGCC-3'

TRANSLATION TRANSCRIPTION
 STOP CODONS STOP SIGNAL
 SALI

TTCGATATCAAGCTTATCGATACCGTCGACCTCGAGGGGGCTAACTAACTAATTGTTTGT

APAI

GGGGCCGGCC-3'

FIG. 2

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3. p7.5/ATG1/tk

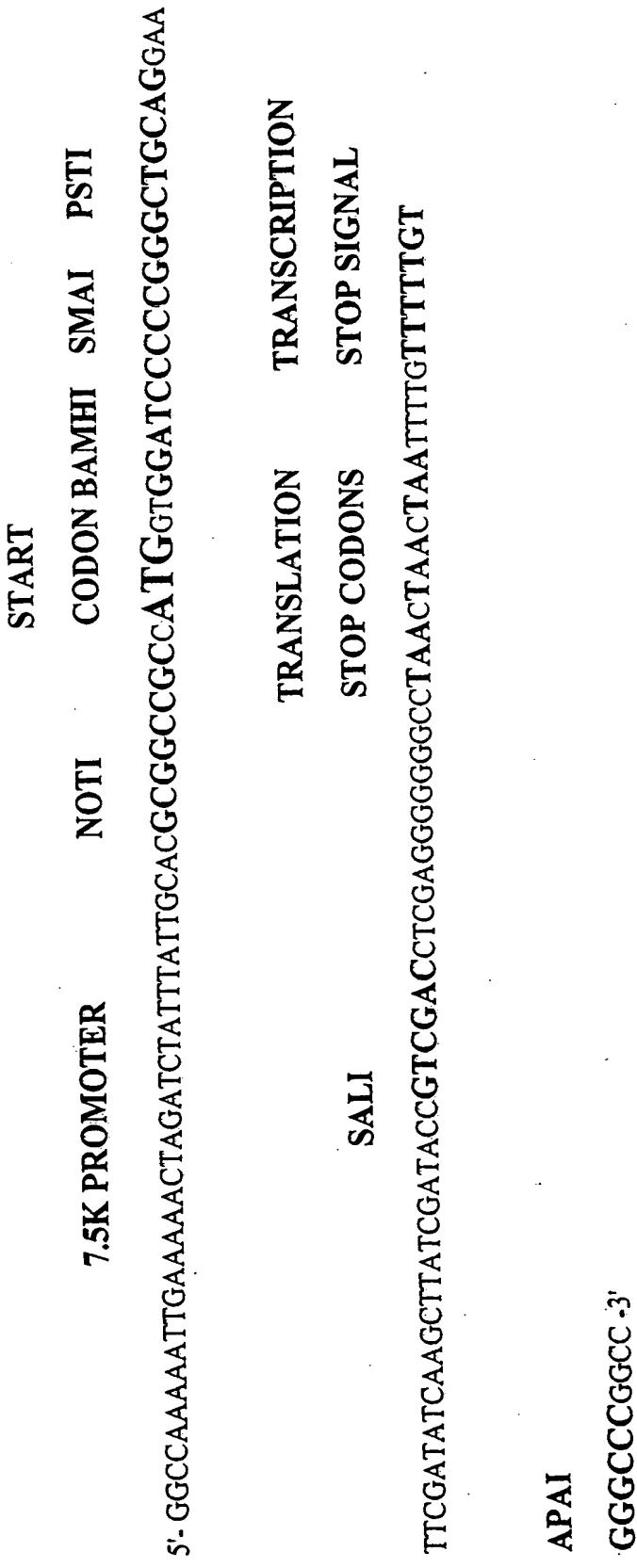
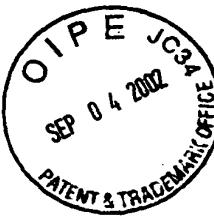


FIG. 2 (cont.)



4. p7.5/ATG2/tk

START

7.5K PROMOTER

NOTI CODON BAMHI SMAI PSTI

5: GGCCAAAAATTGAAACTAGATCTATTGCACGGCCCCCATGAGTGGATCCCCGGCTGCAGGAA

TRANSLATION

STOP CODONS

SALI

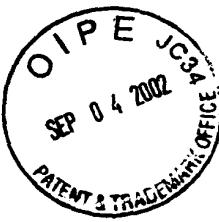
TTCGATATCAAGCTTATCGATAACCGTCGACCTCGAGGGGGGCCTAACTAACTAATTGTTTTGT

APAI

GGGGCCCCGCC-3'

FIG. 2 (cont.)

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5. p7.5/ATG3/tk

7.5K PROMOTER

NOTI

START

CODON BAMHI SMAI PSTI

5'-GGCCAAAAATTGAAAAGTAACTAGATCTATTATTGCACGGGCCATGACGTTGGATCCCCGGCTGCAGGAA

TRANSLATION

TRANSCRIPTION

STOP CODONS

STOP SIGNAL

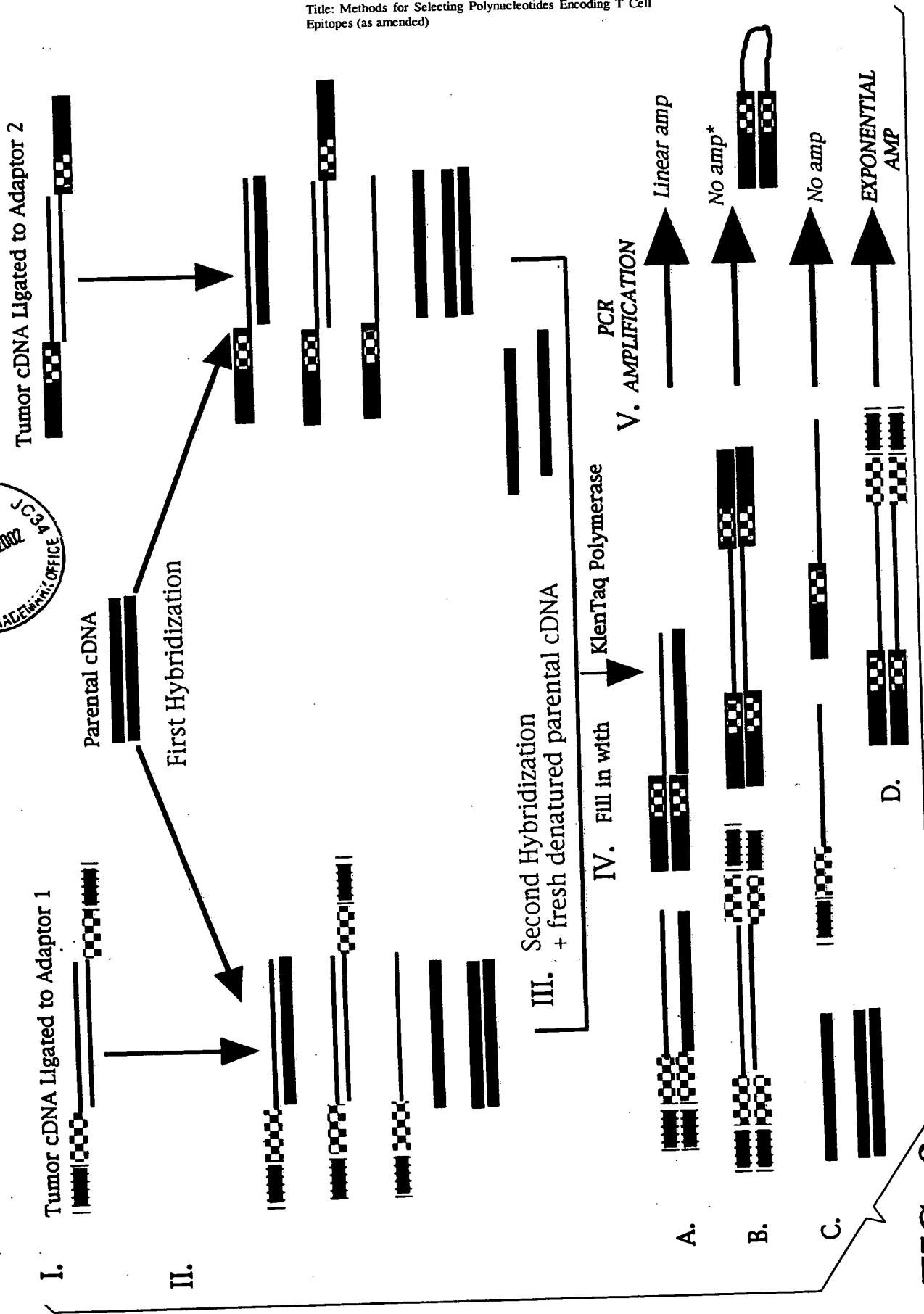
SALI

TTCGATATCAAGCTTATCGATAACCGTCGACCTCGAGGGGGGGCCTAACTAACTAATTGTTTTGT

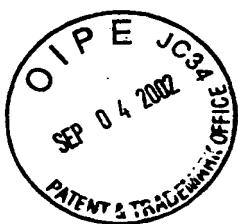
APAI

GGGGGGGGCC-3'

FIG. 2 (cont.)

**FIG. 3**

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SUBTRACTED
100 bp LADDER

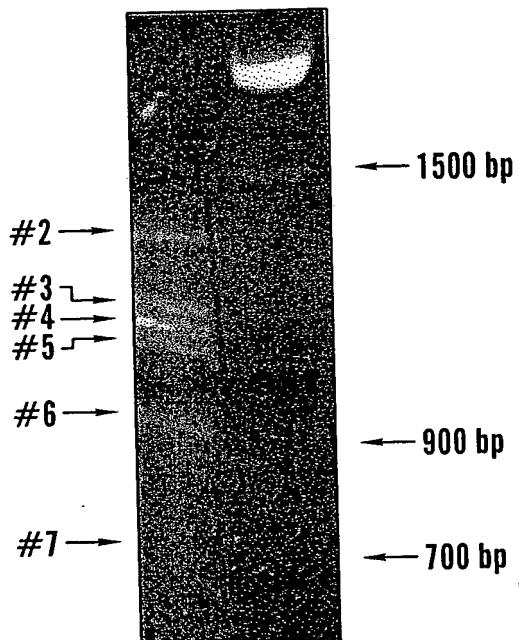


FIG. 4

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G3PDH:

TUMOR

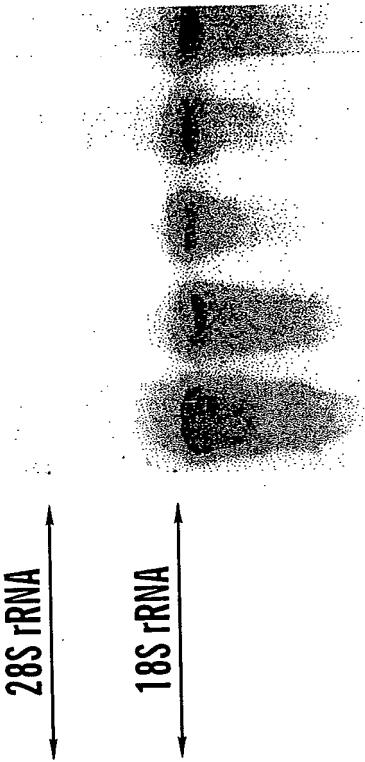
BCA 39

BCA 34

BCA 22

BCB 13

B/cN PARENTAL

CLONE #1:

TUMOR

BCA 39

BCA 34

BCA 22

BCB 13

B/cN PARENTAL

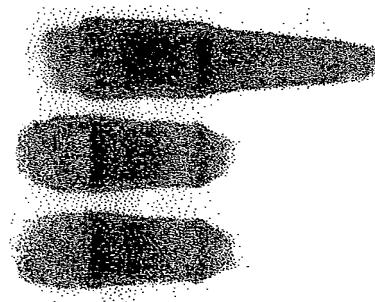
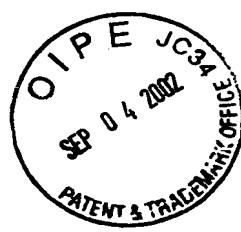


FIG. 5

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- = Region of the insert that has been sequenced
- = Estimated region of IAP gene encoded by the insert, based on the molecular weight of the cloned RDA band
- ↖ = Deletion in IAP sequence, i.e. distance between sequences from 5' and 3' ends of the insert is less than the distance homologous to the full length MIA14 sequence

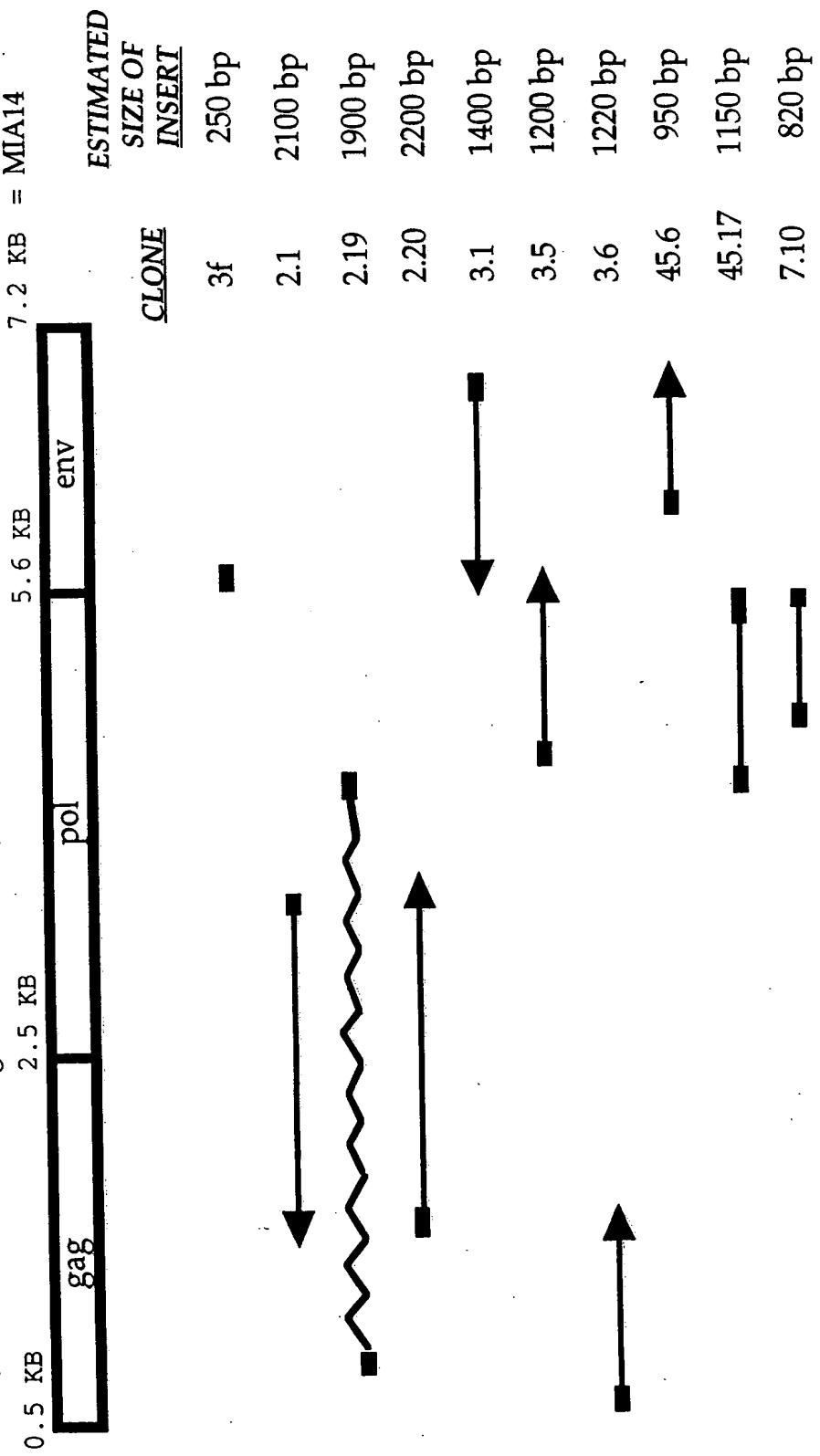


FIG. 6

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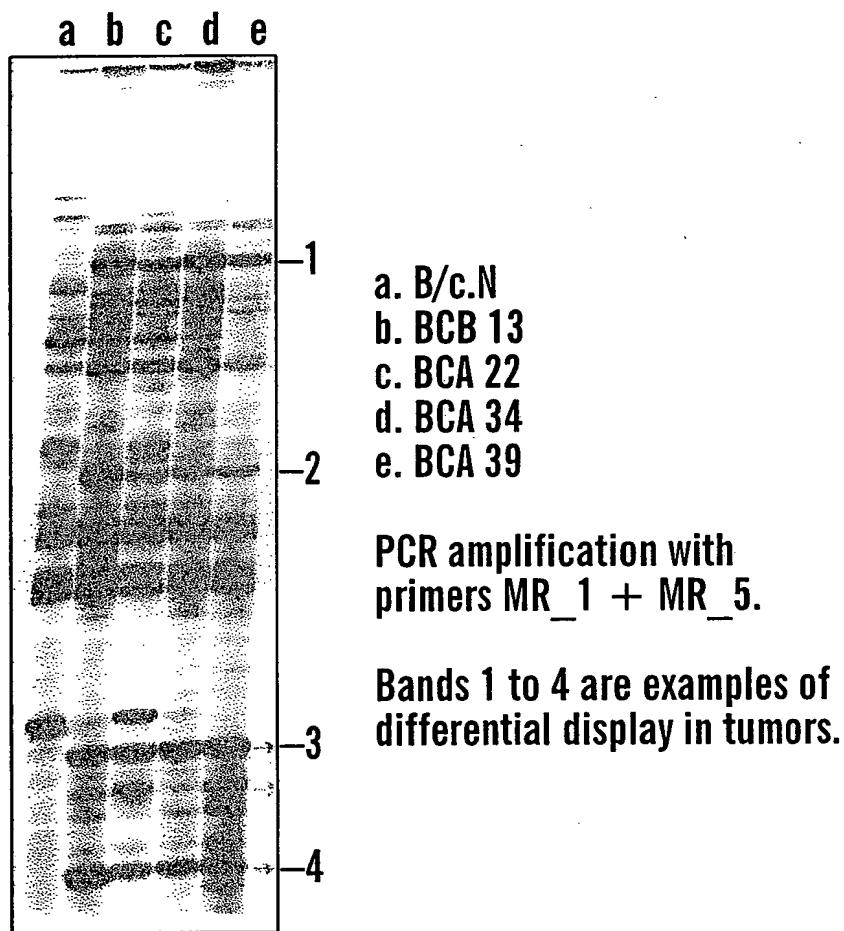
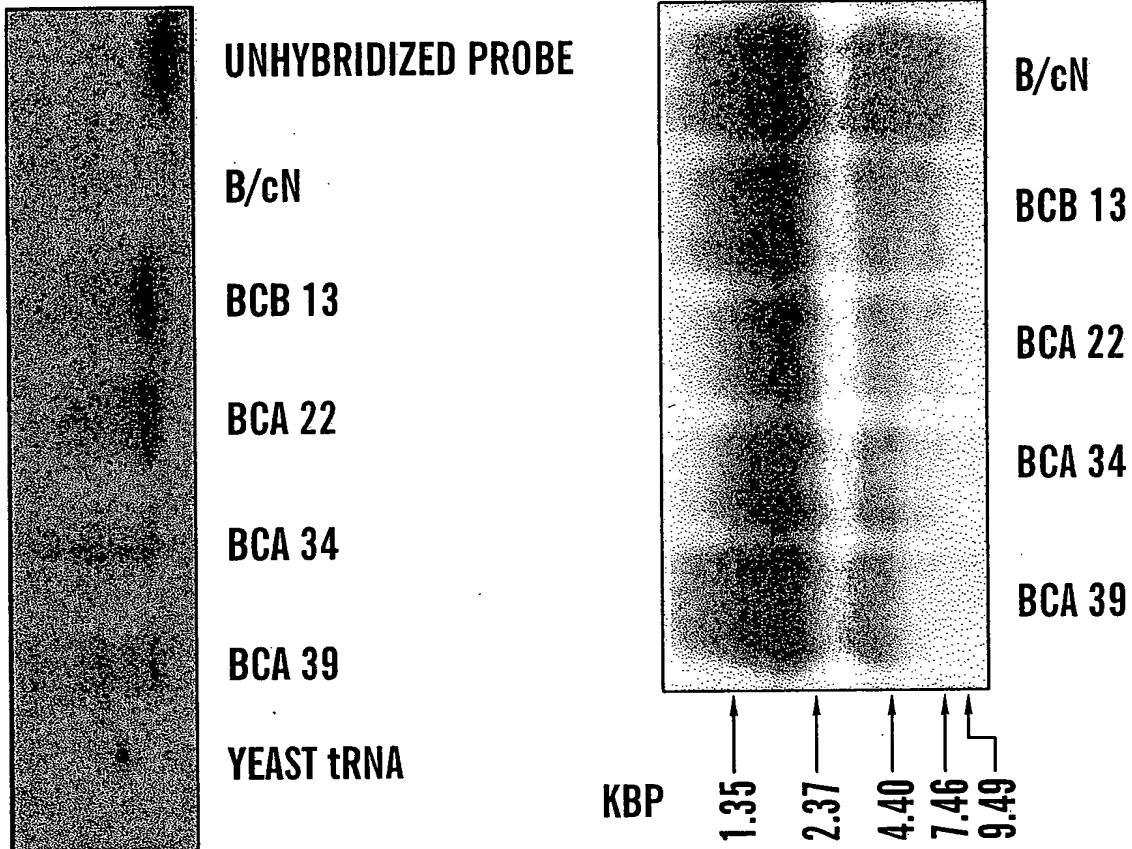
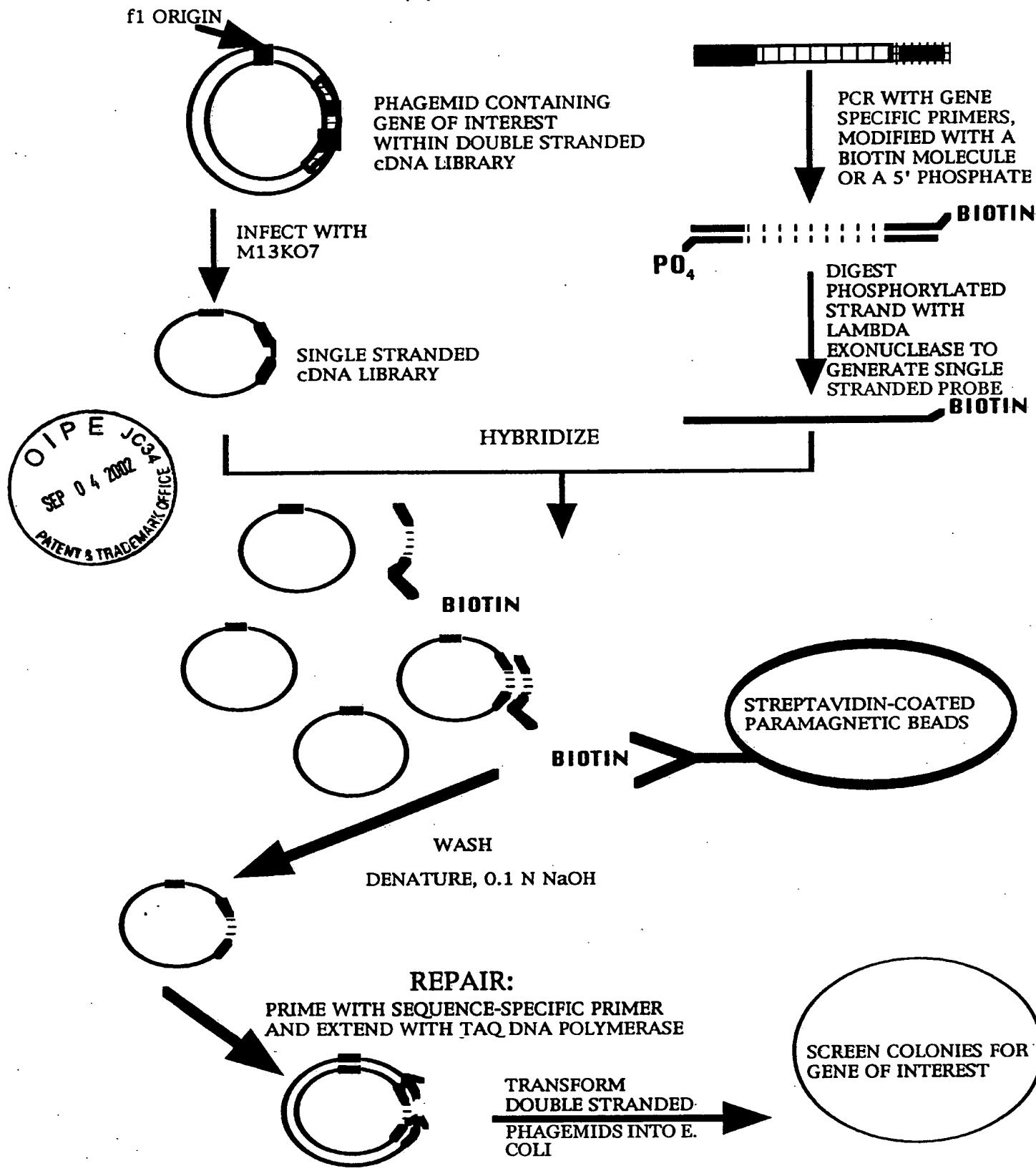


FIG. 7

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**FIG. 9**

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Apal NotI

λ WR WR VEL V7.5 vNotI/tk
WR VEL V7.5 vNotI/tk

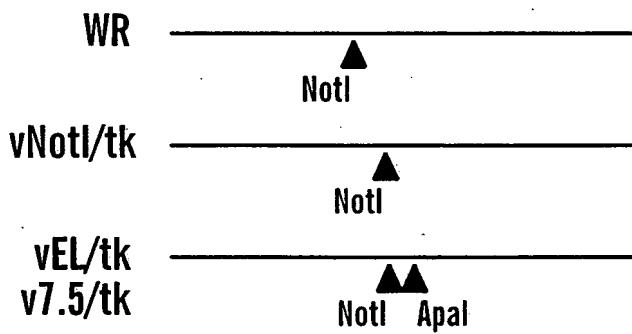
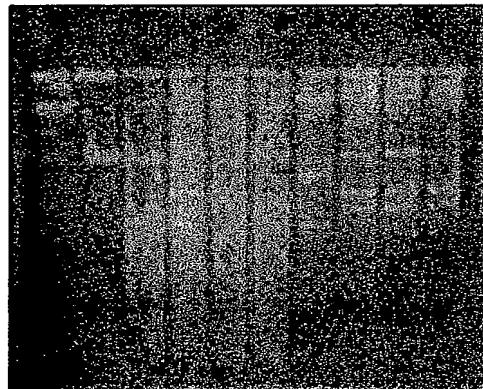


FIG. 10

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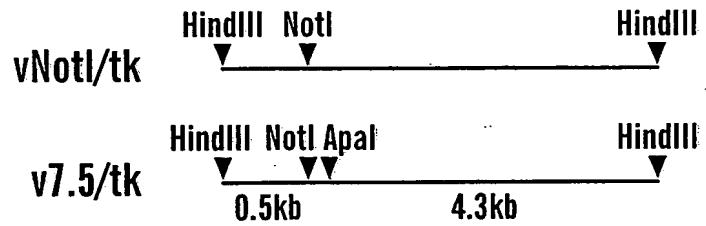
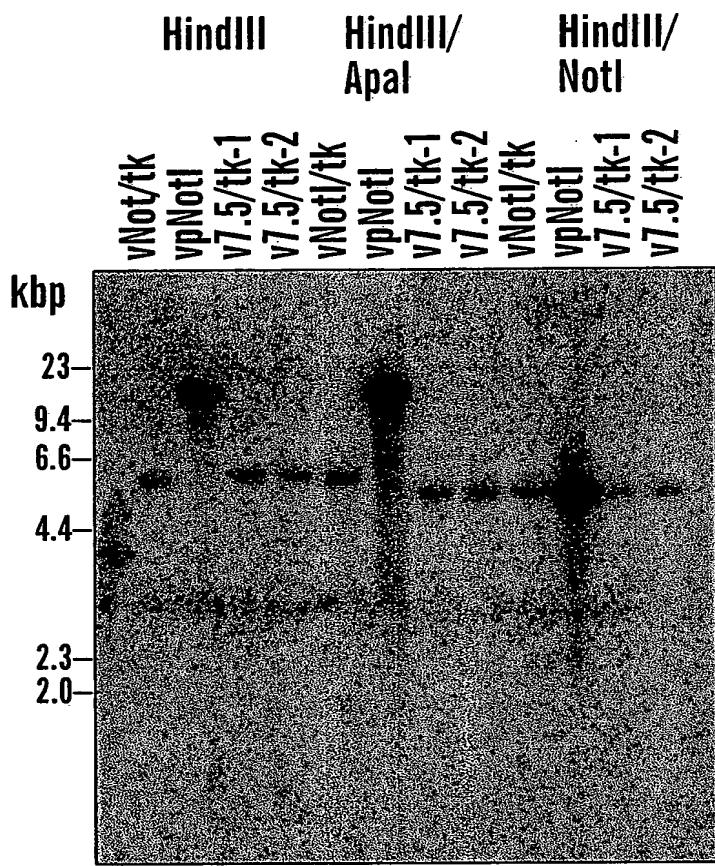


FIG. 11A

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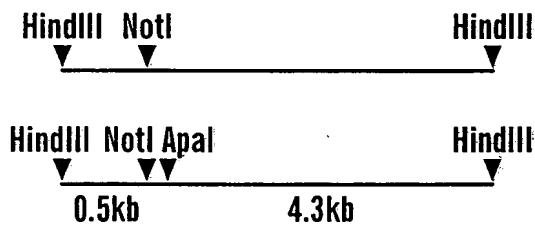
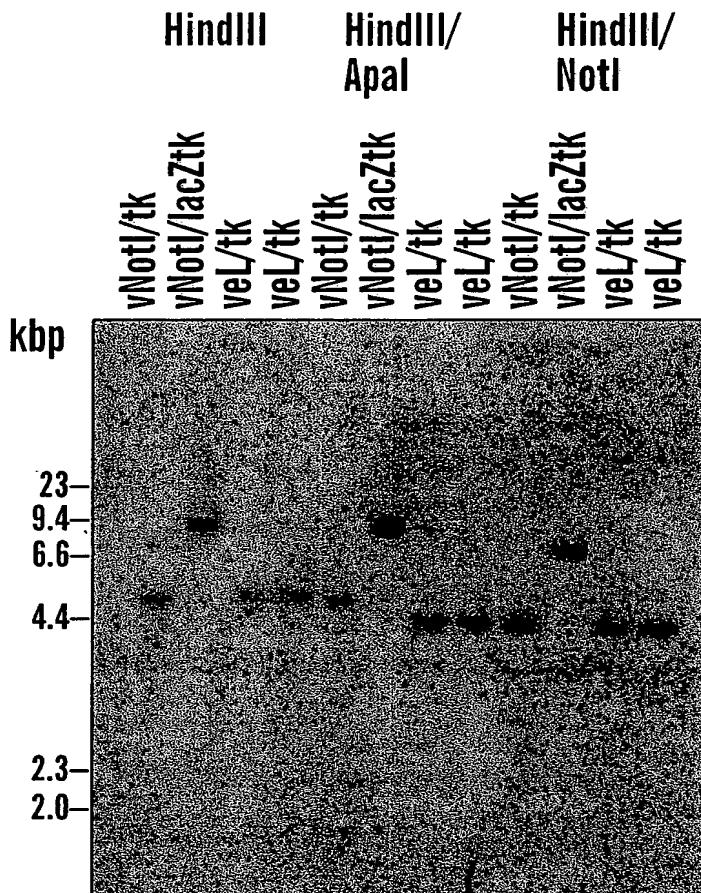


FIG. 11B

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Epitopes (as amended)



Phi X
WR
pNOT
vNotI/tk
vpNotI
p7.5/tk
v7.5/tk
vNotI/lacZ/tk
pel/tk
vel/tk

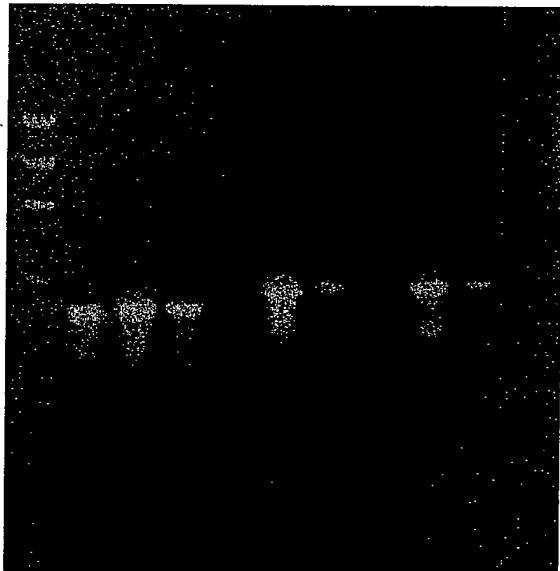


FIG. 12

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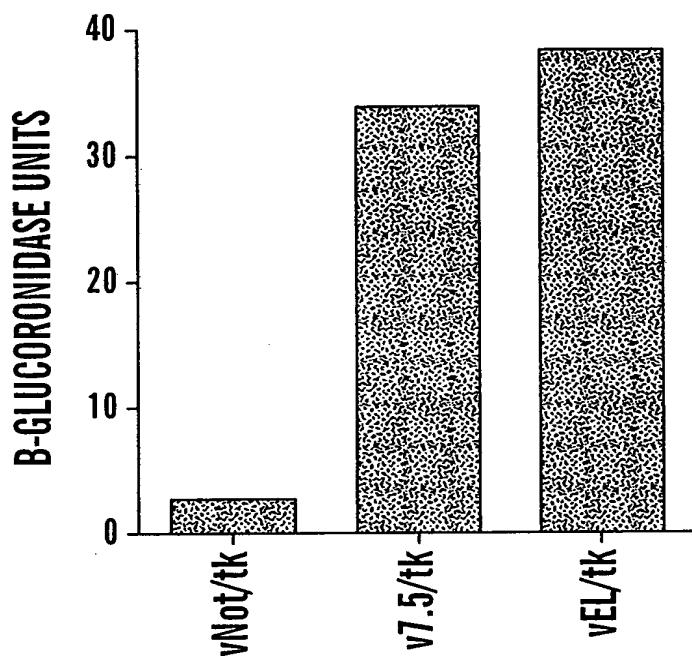


FIG. 13

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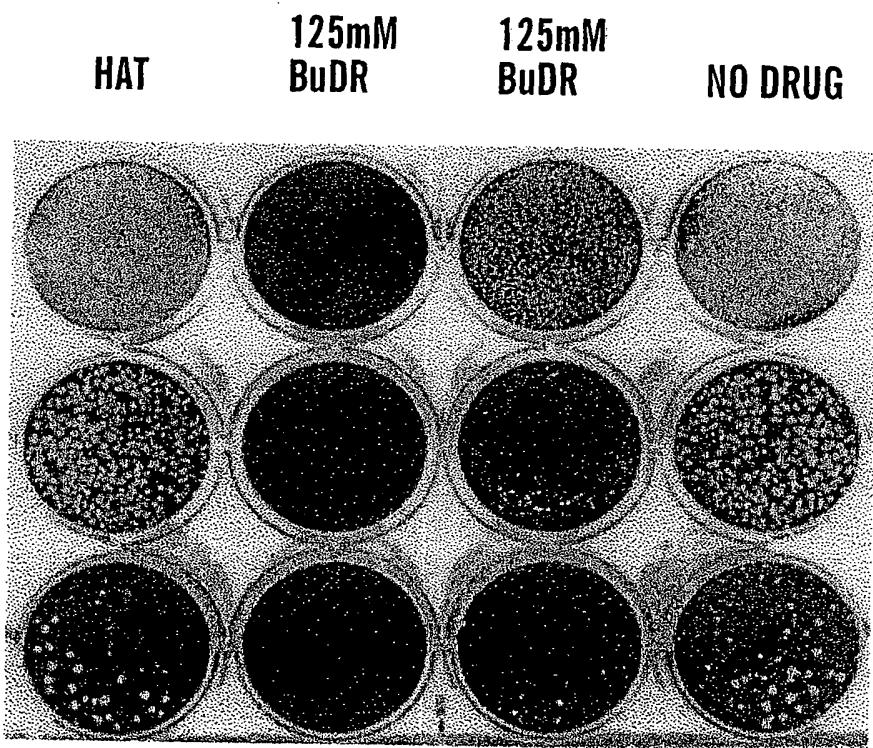


FIG. 14